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=> s saccharomyces (w)cerevisiae L2 423082 SACCHAROMYCES (W) CEREVISIAE

=> s l1 or l2 L3 430360 L1 OR L2

=> s (transform? or transfect?)(w)13
L4 579 (TRANSFORM? OR TRANSFECT?)(W) L3

=> s hybrid (w)sensor(w)kinase? L5 140 HYBRID (W) SENSOR(W) KINASE?

=> d 1-2 ibib ab

L7 ANSWER 1 OF 2 BIOTECHDS COPYRIGHT 2008 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-15129 BIOTECHDS

TITLE: New transformed cell in which a polynucleotide coding for

osmosensing histidine kinase having no

transmembrane region has been introduced, useful for identifying an antifungal compound useful for killing a

fungus;

vector expression in host cell for use in drug screening

and fungus infection therapy

AUTHOR: NAKAJIMA H

PATENT ASSIGNEE: SUMITOMO CHEM CO LTD PATENT INFO: EP 1415996 6 May 2004

APPLICATION INFO: EP 2003-256895 30 Oct 2003 PRIORITY INFO: JP 2002-317736 31 Oct 2002; JP 2002-317736 31 Oct 2002

PRIORITY INFO: JP 2002-317736 31
DOCUMENT TYPE: Patent

DOCUMENT TYPE: Patent LANGUAGE: English

OTHER SOURCE: WPI: 2004-341880 [32]

AB DERWENT ABSTRACT:

NOVELTY - A transformed cell in which a polynucleotide having a sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region has been introduced in a functional form into a cell deficient in at least one hybridsensor kinase, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for: (1) assaying the antifungal activity of a substance; (2) identifying an antifungal compound; (3) an antifungal compound selected by the method above; (4) killing a fungus; (5) an osmosensing histidine kinase having no transmembrane region and derived from a plant-pathogenic filamentous fungus or which has an amino acid sequence selected from: (a) a sequence of 1293, 1307 or 1438 amino acids (SEQ ID NO: 41, 55 or 68) given in the specification or a sequence 95% or more homologous to SEQ ID NO: 41, 55 or 68; (b) a sequence encoded by a DNA amplified by PCR using a Fusarium oxysporum-derived cDNA as a template and using oligonucleotides having the nucleotide sequences tgcactagtatggttgacgacgcggccctcgc (SEQ ID NO: 52) and gagctgcagttagttggtaagacttcgcatatc (SEQ ID NO: 53) as primers; (c) a sequence encoded by a DNA amplified by PCR using Mycospharella tritici-derived cDNA as a template and using oligonucleotides having the sequences cccactagtatgctgcaagaagagacttcg (SEQ ID NO: 64) and cctaagcttctcagctgctatgggcacgaa (SEQ ID NO: 65) as primers; (d) a sequence encoded by a DNA amplified by PCR using Thanapethorus cucumeris-derived cDNA as a template and using oligonucleotides having the sequences ggaactagtatggcaggtacaacggggggacacc (SEQ ID NO: 85) and tgcaagcttttagtgggcaccgtggggtgttacg (SEQ ID NO: 86) as primers; and (e) a sequence derived from Phytophthora infestans and has the amino acid sequence of 124 amino acids (SEQ ID NO: 90) given in the specification; (6) a polynucleotide having a nucleotide sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region derived from a plant-pathogen filamentous fungus described above or having a sequence of 3882, 3924, or 4317 bp (SEQ ID NO: 42, 56, or 69) given in the specification; (7) obtaining the polynucleotide above; and (8) an oligonucleotide which comprises a nucleotide sequence selected from 17 sequences of 23-34 bp (SEQ ID NO: 30-40, 52, 53, 64, 65, 85, and 86) given in the specification, e.g., aacatqtcccacqarattcgmacacc (SEQ ID NO: 30) caccgagattcgvacacccatgaaygg (SEQ ID NO: 31) aggccttccaaaaggctctvcggga (SEQ ID NO: 32) gagatggaccctgaaatcacmac (SEQ ID NO: 33) cagatattctcyagygaagtytckcg (SEQ ID NO: 34) atagcrttgccaacmaggttmagaataa (SEQ ID NO: 35)

aacttgatggcrttkccaacmaggtt (SEQ ID NO: 36) ctctgtgaacttgatrgcrttkccaac (SEQ ID NO: 37) atacacttttcncggtcacccatcat (SEQ ID NO: 38) tccatctgbgcctggatacacttttc (SEQ ID NO: 39) ggcttygavagatactcgtccatctg (SEQ ID NO: 40).

BIOTECHNOLOGY - Preferred Transformed Cell: The polynucleotide is a polynucleotide complementing the deficiency in hybridsensor kinase in the cell deficient in at least one hybrid-sensor kinase in which the polynucleotide has been introduced. The cell is a microorganism, particularly budding yeast. The osmosensing histidine kinase having no transmembrane region is an osmosensing histidine kinase having no transmembrane region and having a mutation which confers resistance to any of a dicarboxylimide antifungal compound, an aromatic hydrocarbon antifungal compound and a phenylpyrrole antifungal compound to the cell. The osmosensing histidine kinase is derived from a plant-pathogenic filamentous fungus and has no transmembrane region. The osmosensing histidine kinase has an amino acid sequence of 1315, 1307, 1293, 1307, 1438 or 124 amino acids (SEQ ID NO: 1, 16, 41, 55, 68 or 90, respectively) given in the specification. The nucleotide sequence encoding an amino acid sequence of the osmosensing histidine kinase is a sequence of 3948, 3924, 3882, 3924, or 4317 bp (SEQ ID NO: 2, 17, 42, 56, or 69, respectively) also given in the specification. Preferred Method: Assaying the antifungal activity of a substance comprises culturing a transformed cell defined above in the presence of a test substance, measuring an amount of intracellular signal transduction from the osmosensing histidine kinase having no transmembrane region expressed in the cultured transformed cell or an index value having the correlation, and assessing the antifungal activity of the test substance based on a difference between an amount of intracellular signal transduction or an index value having the correlation measured and a control. The amount of intracellular signal transduction or the index value having the correlation is an amount of growth of the transformed cell. Identifying an antifungal compound comprises selecting an antifungal compound based on the antifungal activity assessed in the assaying method defined above. Killing a fungus comprises identifying an antifungal compound by the method above and contacting the fungus with the identified antifungal compound. Obtaining the polynucleotide above comprises amplifying a desired polynucleotide by PCR using the oligonucleotide above and recovering the amplified desired polynucleotide.

ACTIVITY - Fungicide. No biological data given. MECHANISM OF ACTION - None given.

USE - The transformed cell is useful for assaying the antifungal activity of a substance and identifying an antifungal compound which is useful for killing a fungus (claimed).

EXAMPLE - BcOS-1 DNA was cloned into a shuttle vector p415ADH (ATCC87312) replicable in yeast and Escherichia coli. Both were digested, separated by agarose gel electrophoresis, and a part of the gel containing a desired DNA was excised. The BcOS-1 DNA was inserted between SpeI and Pstl sites in the multicloning site of the shuttle vector. A nucleotide sequence of the resulting expression plasmid was analyzed after a sequencing reaction. The nucleotide sequence of 3948 bp (SEQ ID NO: 2) given in the specification was obtained and it was confirmed that the expression plasmid pADHBcOS1 harbored a DNA having a nucleotide sequence encoding an amino acid sequence of BcOS-1. The prepared expression plasmid was introduced into each of budding yeast (Saccharomyces cerevisiae AH22 strain) (IFO10144) and TM182 strain. By utilizing disappearance of leucine auxotrophy in the resulting transformed budding yeast, the transformed budding yeast AH22 strain was selected on a Glu-Leu agar medium, and the transformed budding yeast TM182 was selected on a Gal-Ura-Leu agar medium. It was confirmed that the resulting TM182-BcOS1 grows even when transplanted to a Glu-Ura-Leu medium. (211 pages)

L7 ANSWER 2 OF 2 HCAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2004:370684 HCAPLUS

DOCUMENT NUMBER: 140:369919

TITLE: Transformed cell with enhanced sensitivity to

antifungal compound, expressing mutated gene, os-1,

for an osmosensing histidine kinase, and uses for fungicide screening

INVENTOR(S): Nakajima, Hiroki

PATENT ASSIGNEE(S): Sumitomo Chemical Company, Limited, Japan

SOURCE: Eur. Pat. Appl., 211 pp.

CODEN: EPXXDW

DOCUMENT TYPE:

Patent English

LANGUAGE: English

FAMILY ACC. NUM. COUNT: PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
EP 1415996	A2	20040506	EP 2003-256895	20031030
EP 1415996	A3	20040901		
EP 1415996	B1	20071017		
R: AT, BE, CH,	DE, DK	, ES, FR, G	B, GR, IT, LI, LU,	NL, SE, MC, PT,
IE, SI, LT,	LV, FI	, RO, MK, C	Y, AL, TR, BG, CZ,	EE, HU, SK
JP 2005087182	Α	20050407	JP 2003-354761	20031015
SG 127705	A1	20061229	SG 2003-6525	20031030
AT 375997	T	20071115	AT 2003-256895	20031030
US 2004137594	A1	20040715	US 2003-697036	20031031
PRIORITY APPLN. INFO.:			JP 2002-317736	A 20021031
			JP 2003-207458	A 20030813

An object of the present invention is to provide a method of detecting the antifungal activity and a method of antifungal screening using filamentous fungi homologs of Neurispora crassa os-1 gene encoding a two-component system osmosensing histidine kinase having no transmembrane region. OS-1 protein and cDNA sequences from phytopathogenic fungi, including Botryotinia fuckeliana (BcOS-1), Magnaoirthe grisea (HIK1), Fusarium oxysporum (FoOS-1), Mycosphaerella tritici (StOS-1), Thanatephorus cucumeris (RsOS-1), and Phytophthora infestans (PiOS-1), are provided. The present invention provides transformed cells (such as budding yeast) in which a os-1 gene homolog encoding an osmosensing histidine kinase having no transmembrane region has been introduced in a functional form into a cell deficient in at least one hybrid-sensor kinase

The os-1 transgene is carrying a mutation which confers resistance to the cell to any of a dicarboxyimide antifungal compound, an aromatic hydrocarbon antifungal compound and a phenylpyrrole antifungal compound Provided are a method of assaying the antifungal activity of a test substance using the transformed cell, and a method of identifying an antifungal compound

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L1 38279 S BUDDING (W) YEAST?

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L3 430360 S L1 OR L2

L4 579 S (TRANSFORM? OR TRANSFECT?) (W) L3

L5 140 S HYBRID (W) SENSOR (W) KINASE?

L6 98 S OSMOSENSING (2W) KINASE?

L7 2 S L5 AND L6

=> d 1-2 ibib ab

L8 ANSWER 1 OF 2 BIOTECHDS COPYRIGHT 2008 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-15129 BIOTECHDS

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and fungus infection therapy

AUTHOR: NAKAJIMA H

PATENT ASSIGNEE: SUMITOMO CHEM CO LTD PATENT INFO: EP 1415996 6 May 2004 APPLICATION INFO: EP 2003-256895 30 Oct 2003

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DOCUMENT TYPE: Patent LANGUAGE: English

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transplanted to a Glu-Ura-Leu medium. (211 pages) ANSWER 2 OF 2 HCAPLUS COPYRIGHT 2008 ACS on STN 2004:370684 HCAPLUS ACCESSION NUMBER: 140:369919 DOCUMENT NUMBER: Transformed cell with enhanced sensitivity to TITLE: antifungal compound, expressing mutated gene, os-1, for an osmosensing histidine kinase , and uses for fungicide screening Nakajima, Hiroki INVENTOR(S): Sumitomo Chemical Company, Limited, Japan PATENT ASSIGNEE(S): Eur. Pat. Appl., 211 pp. SOURCE: CODEN: EPXXDW DOCUMENT TYPE: Patent LANGUAGE: English FAMILY ACC. NUM. COUNT: 1 PATENT INFORMATION: PATENT NO. KIND DATE APPLICATION NO. DATE ----------_____ -----20040506 EP 2003-256895 EP 1415996 A2 20031030 A3 EP 1415996 20040901 B1 20071017 EP 1415996 R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL, TR, BG, CZ, EE, HU, SK A JP 2005087182 20050407 JP 2003-354761 20031015 SG 127705 A1 20061229 SG 2003-6525 20031030 20031030 20071115 AT 2003-256895 AT 375997 T US 2003-697036 A1 US 2004137594 20040715 20031031 A 20021031 A 20030813 JP 2002-317736 PRIORITY APPLN. INFO.: JP 2003-207458

AB An object of the present invention is to provide a method of detecting the antifungal activity and a method of antifungal screening using filamentous fungi homologs of Neurispora crassa os-1 gene encoding a two-component system osmosensing histidine kinase having no transmembrane region. OS-1 protein and cDNA sequences from phytopathogenic fungi, including Botryotinia fuckeliana (BcOS-1), Magnaoirthe grisea (HIK1), Fusarium oxysporum (FoOS-1), Mycosphaerella tritici (StOS-1), Thanatephorus cucumeris (RsOS-1), and Phytophthora infestans (PiOS-1), are provided. The present invention provides transformed cells (such as budding yeast) in which a os-1 gene homolog encoding an osmosensing histidine kinase having no transmembrane region has been introduced in a functional form into a cell

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. The os-1 transgene is carrying a mutation which confers resistance to the cell to any of a dicarboxyimide antifungal compound, an aromatic hydrocarbon antifungal compound and a phenylpyrrole antifungal compound Provided are a method of assaying the antifungal activity of a test substance using the transformed cell, and a method of identifying an antifungal compound

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=> e nakajima h/au
                          NAKAJIMA GORO/AU
                5
E2
                          NAKAJIMA GOZO/AU
                 1
E3
             9245 --> NAKAJIMA H/AU
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14 NAKAJIMA HACHIRO/AU
14 NAKAJIMA HADJIME/AU
152 NAKAJIMA HAJIME/AU
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L9 9245 "NAKAJIMA H"/AU
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- FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 09:28:41 ON 18 JAN 2008

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L8 2 S L4 AND (L5 OR L6) E NAKAJIMA H/AU

L9 9245 S E3

=> s 14 and 19

L10 1 L4 AND L9

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L10 ANSWER 1 OF 1 BIOTECHDS COPYRIGHT 2008 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-15129 BIOTECHDS

TITLE:

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vector expression in host cell for use in drug screening

and fungus infection therapy

AUTHOR: NAKAJIMA H

PATENT ASSIGNEE: SUMITOMO CHEM CO LTD
PATENT INFO: EP 1415996 6 May 2004
APPLICATION INFO: EP 2003-256895 30 Oct 2003

PRIORITY INFO: JP 2002

JP 2002-317736 31 Oct 2002; JP 2002-317736 31 Oct 2002

DOCUMENT TYPE: Patent LANGUAGE: English

OTHER SOURCE: WPI: 2004-341880 [32]

AB DERWENT ABSTRACT:

NOVELTY - A transformed cell in which a polynucleotide having a sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region has been introduced in a functional form into a cell deficient in at least one hybrid-sensor kinase, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for: (1) assaying the antifungal activity of a substance; (2) identifying an antifungal compound; (3) an antifungal compound selected by the method above; (4) killing a fungus; (5) an osmosensing histidine kinase having no transmembrane region and derived from a plant-pathogenic filamentous fungus or which has an amino acid sequence selected from: (a) a sequence of 1293, 1307 or 1438 amino acids (SEQ ID NO: 41, 55 or 68) given in the specification or a sequence 95% or more homologous to SEQ ID NO: 41, 55 or 68; (b) a sequence encoded by a DNA amplified by PCR using a Fusarium oxysporum-derived cDNA as a template and using oligonucleotides having the nucleotide sequences tgcactagtatgqttgacgacgcggccctcgc (SEQ ID NO: 52) and qaqctqcaqttaqttqqtaaqacttcqcatatc (SEQ ID NO: 53) as primers; (c) a sequence encoded by a DNA amplified by PCR using Mycospharella tritici-derived cDNA as a template and using oligonucleotides having the sequences cccactagtatgctgcaagaagagacttcg (SEQ ID NO: 64) and cctaagcttctcagctgctatgggcacgaa (SEQ ID NO: 65) as primers; (d) a sequence encoded by a DNA amplified by PCR using Thanapethorus cucumeris-derived cDNA as a template and using oligonucleotides having the sequences ggaactagtatggcaggtacaacggggggacacc (SEQ ID NO: 85) and tgcaagcttttagtgggcaccgtggggtgttacg (SEQ ID NO: 86) as primers; and (e) a sequence derived from Phytophthora infestans and has the amino acid sequence of 124 amino acids (SEQ ID NO: 90) given in the specification; (6) a polynucleotide having a nucleotide sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region derived from a plant-pathogen filamentous fungus described above or having a sequence of 3882, 3924, or 4317 bp (SEQ ID NO: 42, 56, or 69) given in the specification; (7) obtaining the polynucleotide above; and (8) an oligonucleotide which comprises a nucleotide sequence selected from 17 sequences of 23-34 bp (SEQ ID NO: 30-40, 52, 53, 64, 65, 85, and 86) given in the specification, e.g., aacatgtcccacgarattcgmacacc (SEQ ID NO: 30) caccgagattcgvacacccatgaaygg (SEQ ID NO: 31) aggeetteeaaaaggetetveggga (SEQ ID NO: 32) gagatggaceetgaaateacmae (SEQ ID NO: 33) cagatattctcyagygaagtytckcg (SEQ ID NO: 34) atagcrttgccaacmaggttmagaataa (SEQ ID NO: 35) aacttgatggcrttkccaacmaggtt (SEQ ID NO: 36) ctctgtgaacttgatrgcrttkccaac (SEQ ID NO: 37) atacacttttcncggtcacccatcat (SEQ ID NO: 38) tccatctgbgcctggatacacttttc (SEQ ID NO: 39) ggcttvgavagatactcgtccatctg (SEQ ID NO: 40).

BIOTECHNOLOGY - Preferred Transformed Cell: The polynucleotide is a polynucleotide complementing the deficiency in hybrid-sensor kinase in the cell deficient in at least one hybrid-sensor kinase in which the polynucleotide has been introduced. The cell is a microorganism, particularly budding yeast. The osmosensing histidine kinase having no transmembrane region is an osmosensing histidine kinase having no transmembrane region and having a mutation which confers resistance to any of a dicarboxylimide antifungal compound, an aromatic hydrocarbon antifungal compound and a phenylpyrrole antifungal compound to the cell. The osmosensing histidine kinase is derived from a plant-pathogenic filamentous fungus and has no transmembrane region. The osmosensing histidine kinase has an amino acid sequence of 1315, 1307, 1293, 1307, 1438 or 124 amino acids (SEQ ID NO: 1, 16, 41, 55, 68 or 90, respectively) given in the specification. The nucleotide sequence encoding an amino acid sequence of the osmosensing histidine kinase is a sequence of 3948, 3924, 3882, 3924, or 4317 bp (SEQ ID NO: 2, 17, 42, 56, or 69, respectively) also given in the specification. Preferred Method: Assaying the antifungal activity of a substance comprises culturing a transformed cell defined above in the presence of a test substance, measuring an amount of intracellular signal transduction from the osmosensing histidine kinase having no transmembrane region expressed in the cultured transformed cell or an index value having the correlation, and assessing the antifungal activity of the test substance based on a difference between an amount of intracellular signal transduction or an index value having the correlation measured and a control. The amount of intracellular signal transduction or the index value having the correlation is an amount of growth of the transformed cell. Identifying an antifungal compound comprises selecting an antifungal compound based on the antifungal activity assessed in the assaying method defined above. Killing a fungus comprises identifying an antifungal compound by the method above and contacting the fungus with the identified antifungal compound. Obtaining the polynucleotide above comprises amplifying a desired polynucleotide by PCR using the oligonucleotide above and recovering the amplified desired polynucleotide.

ACTIVITY - Fungicide. No biological data given. MECHANISM OF ACTION - None given.

USE - The transformed cell is useful for assaying the antifungal activity of a substance and identifying an antifungal compound which is useful for killing a fungus (claimed).

EXAMPLE - BcOS-1 DNA was cloned into a shuttle vector p415ADH (ATCC87312) replicable in yeast and Escherichia coli. Both were digested, separated by agarose gel electrophoresis, and a part of the gel containing a desired DNA was excised. The BcOS-1 DNA was inserted between

SpeI and Pstl sites in the multicloning site of the shuttle vector. A nucleotide sequence of the resulting expression plasmid was analyzed after a sequencing reaction. The nucleotide sequence of 3948 bp (SEQ ID NO: 2) given in the specification was obtained and it was confirmed that the expression plasmid pADHBcOS1 harbored a DNA having a nucleotide sequence encoding an amino acid sequence of BcOS-1. The prepared expression plasmid was introduced into each of budding yeast (Saccharomyces cerevisiae AH22 strain) (IFO10144) and TM182 strain. By utilizing disappearance of leucine auxotrophy in the resulting transformed budding yeast, the transformed budding yeast AH22 strain was selected on a Glu-Leu agar medium, and the transformed budding yeast TM182 was selected on a Gal-Ura-Leu agar medium. It was confirmed that the resulting TM182-BcOS1 grows even when transplanted to a Glu-Ura-Leu medium. (211 pages)

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L10

(FILE 'HOME' ENTERED AT 09:28:11 ON 18 JAN 2008)

1 S L4 AND L9

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FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS,
     LIFESCI' ENTERED AT 09:28:41 ON 18 JAN 2008
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L2
         430360 S L1 OR L2
L3
            579 S (TRANSFORM? OR TRANSFECT?) (W) L3
L4
            140 S HYBRID (W) SENSOR (W) KINASE?
L5
             98 S OSMOSENSING (2W) KINASE?
L6
              2 S L5 AND L6
L7
              2 S L4 AND (L5 OR L6)
L8
                E NAKAJIMA H/AU
           9245 S E3
L9
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L12 ANSWER 1 OF 4 HCAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2006:1145495 HCAPLUS

TITLE: Signal transduction in yeast involving a His-to-Asp

phosphorelay system

AUTHOR(S): West, Ann H.; Xu, Qingping; Porter, Stace;

Janiak-Spens, Fabiola; Chooback, Lilian

CORPORATE SOURCE: Department of Chemistry and Biochemistry, University

of Oklahoma, Norman, OK, 73019, USA

SOURCE: Abstracts, 62nd Southwest Regional Meeting of the

American Chemical Society, Houston, TX, United States,

October 19-22 (2006), SRM-131. American Chemical

Society: Washington, D. C.

CODEN: 69INSW

DOCUMENT TYPE: Conference; Meeting Abstract

LANGUAGE: English

Histidine-containing phosphotransfer (HPt) proteins play an essential role in the transfer of phosphoryl groups between response regulator domains in multi-step His-Asp signal transduction systems. In Saccharomyces cerevisiae, the HPt protein YPD1 facilitates phosphoryl group transfer from the response regulator domain of a membrane-bound hybrid sensor kinase (SLN1) to two downstream response regulators SSK1 and SKN7, which are involved in hyperosmotic and oxidative stress responses, resp. Protein-protein interactions involving signaling partners are transient and phosphorylation-dependent. This talk will focus on the mol. basis of interaction of YPD1 with each of the three response regulator domains (SLN1-R1, SSK1-R2, and SKN7-R3, resp.), which we have investigated through a variety of means, including a comparative yeast two-hybrid interaction assay, X-ray crystallog. anal. of a YPD1/SLN1-R1 complex, and kinetic anal. of YPD1-dependent phosphotransfer

L12 ANSWER 2 OF 4 BIOTECHDS COPYRIGHT 2008 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2005-14092 BIOTECHDS

TITLE: Novel transformed cell produced by transducing gene encoding

osmotic-pressure sensitive histidine kinase that functions on

cell lacking hybrid sensor kinase

function, useful for screening antimicrobial agents;

yeast host cell transformation using fungus enzyme gene

for use in antimicrobial substance screening

PATENT ASSIGNEE: SUMITOMO CHEM CO LTD PATENT INFO: JP 2005087182 7 Apr 2005 APPLICATION INFO: JP 2003-354761 15 Oct 2003

PRIORITY INFO: JP 2003-207458 13 Aug 2003; JP 2002-317736 31 Oct 2002

DOCUMENT TYPE: Patent LANGUAGE: Japanese

OTHER SOURCE: WPI: 2005-288601 [30]

AB DERWENT ABSTRACT:

reactions.

NOVELTY - A transformed cell (I), produced by transducing a gene having a base sequence encoding osmotic-pressure sensitive histidine kinase that does not have cytoplasmic membrane penetration region that functions on the cell, where the cell lacks the hybrid sensor kinase function, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are included for the following: (1) testing (M1) antimicrobial activity of a substance; (2) screening (M2) anti-microbial substance; (3) anti-microbial substance (II) screened by (M2); (4) pathogenic, filamentous plant fungi (III), comprising osmotic pressure sensitive histidine kinase that does not have cytoplasmic membrane penetration region, or a polynucleotide encoding osmotic pressure sensitive histidine kinase lacking cytoplasmic membrane penetration region; (5) osmotic pressure sensitive histidine kinase lacking cytoplasmic membrane penetration region (IV), comprising a fully defined 1293 amino acid (SEQ ID No:41), 1307 amino acid (SEQ ID No:55), 1438 amino acid (SEQ ID No:68) sequences given in the specification, one

or more addition, deletion or substitution in SEQ ID No:41, 55 and 69, and SEQ ID No:41 and SEQ ID No: 55 having 95% or more sequence identity with SEQ ID No:68, or a base sequence encoding SEQ ID No:41, 55 and 68; (6) polynucleotide (V) having a fully defined 3882 nucleotide (SEQ ID No:42), 3924 nucleotide (SEQ ID No:56) and 4317 nucleotide (SEQ ID No:69) sequences given in the specification; (7) acquisition method of a polynucleotide encoding osmotic pressure sensitive histidine kinase lacking cytoplasmic membrane penetration region; and (8) oligonucleotide comprising a fully defined 10 nucleotide sequence having approximately 23-28 nucleotide (SEQ ID No:30-40) sequences given in the specification, a fully defined 32 nucleotide (SEQ ID No:52), 33 nucleotide (SEQ ID No:53), 30 nucleotide (SEQ ID No:64), 30 nucleotide (SEQ ID No:65), 34 nucleotide (SEQ ID No:85) and 34 nucleotide (SEQ ID No:86) sequences given in the specification.

BIOTECHNOLOGY - Preferred Method: In (I), the gene encoding osmotic-pressure sensitive histidine kinase lacking cytoplasmic membrane penetration region, complements the function of deleted hybrid sensor kinase within the cell lacking hybrid sensor kinase. The transformed cell is a microorganism, preferably budding yeast. The osmotic-pressure sensitive histidine kinase has resistance with respect to dicarboxyimide, aromatic-hydrocarbon or phenyl-pyrrole antimicrobial substance. The osmotic-pressure sensitive histidine kinase is of plant pathogenic mold origin, preferably botrytis disease filamentous fungi, rice-blight filamentous fungi, spinach wilt-disease filamentous fungi, wheat leaf-blight filamentous fungi, rice sheath-blight-disease filament form, or tomato late blight filamentous-fungi origin, and lacks cytoplasmic-membrane penetration region. The osmotic-pressure sensitive histidine kinase has a fully defined 1315 amino acid (SEQ ID No:1), 1307 amino acid (SEQ ID No:16), 1293 amino acid (SEQ ID No:41), 1307 amino acids (SEQ ID No:55) and 1438 amino acids (SEQ ID No:68) sequences given in the specification. The base sequence encoding osmotic-pressure sensitive histidine kinase, has a fully defined 3948 nucleotide (SEQ ID No:2), 3924 nucleotide (SEQ ID No:17), 3882 nucleotide (SEQ ID No:42), 3924 nucleotide (SEQ ID No:56), 4317 nucleotide (SEQ ID No:69) sequences given in the specification.

ACTIVITY - Antimicrobial. No biological data is given.

MECHANISM OF ACTION - Osmotic-pressure sensitive histidine kinase inhibitor.

USE - (I) is useful for screening antimicrobial substances (claimed).

ADVANTAGE - The sensitivity to antimicrobial substance, is increased. (54 pages)

ANSWER 3 OF 4 BIOTECHDS COPYRIGHT 2008 THE THOMSON CORP. on STN DUPLICATE 1

ACCESSION NUMBER: 2004-15129 BIOTECHDS

TITLE:

New transformed cell in which a polynucleotide coding for osmosensing histidine kinase having no transmembrane region has been introduced, useful for identifying an antifungal compound useful for killing a fungus;

vector expression in host cell for use in drug screening and fungus infection therapy

NAKAJIMA H

AUTHOR: SUMITOMO CHEM CO LTD PATENT ASSIGNEE: PATENT INFO: EP 1415996 6 May 2004

APPLICATION INFO: EP 2003-256895 30 Oct 2003

JP 2002-317736 31 Oct 2002; JP 2002-317736 31 Oct 2002 PRIORITY INFO:

DOCUMENT TYPE: Patent English LANGUAGE:

WPI: 2004-341880 [32] OTHER SOURCE:

DERWENT ABSTRACT: AB

NOVELTY - A transformed cell in which a polynucleotide having a sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region has been introduced in a functional form into a cell deficient in at least one hybrid-sensor kinase, is new.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for: (1) assaying the antifungal activity of a substance; (2) identifying an antifungal compound; (3) an antifungal compound selected by the method above; (4) killing a fungus; (5) an osmosensing histidine kinase having no transmembrane region and derived from a plant-pathogenic filamentous fungus or which has an amino acid sequence selected from: (a) a sequence of 1293, 1307 or 1438 amino acids (SEQ ID NO: 41, 55 or 68) given in the specification or a sequence 95% or more homologous to SEQ ID NO: 41, 55 or 68; (b) a sequence encoded by a DNA amplified by PCR using a Fusarium oxysporum-derived cDNA as a template and using oligonucleotides having the nucleotide sequences tgcactagtatggttgacgacgcggccctcgc (SEQ ID NO: 52) and gagetgeagttagttagtaagaettegeatate (SEQ ID NO: 53) as primers; (c) a sequence encoded by a DNA amplified by PCR using Mycospharella tritici-derived cDNA as a template and using oligonucleotides having the sequences cccactaqtatqctqcaaqaagagacttcg (SEQ ID NO: 64) and cctaagcttctcagctgctatgggcacgaa (SEQ ID NO: 65) as primers; (d) a sequence encoded by a DNA amplified by PCR using Thanapethorus cucumeris-derived cDNA as a template and using oligonucleotides having the sequences ggaactagtatggcaggtacaacggggggacacc (SEQ ID NO: 85) and tgcaagcttttagtgggcaccgtggggtgttacg (SEQ ID NO: 86) as primers; and (e) a sequence derived from Phytophthora infestans and has the amino acid sequence of 124 amino acids (SEQ ID NO: 90) given in the specification; (6) a polynucleotide having a nucleotide sequence encoding an amino acid sequence of an osmosensing histidine kinase having no transmembrane region derived from a plant-pathogen filamentous fungus described above or having a sequence of 3882, 3924, or 4317 bp (SEQ ID NO: 42, 56, or 69) qiven in the specification; (7) obtaining the polynucleotide above; and (8) an oligonucleotide which comprises a nucleotide sequence selected from 17 sequences of 23-34 bp (SEQ ID NO: 30-40, 52, 53, 64, 65, 85, and 86) given in the specification, e.g., aacatgtcccacgarattcgmacacc (SEQ ID NO: 30) caccgagattcgvacacccatgaaygg (SEQ ID NO: 31) aggccttccaaaaggctctvcggga (SEQ ID NO: 32) gagatggaccctgaaatcacmac (SEQ ID NO: 33) cagatattctcyagygaagtytckcg (SEQ ID NO: 34) atagcrttgccaacmaggttmagaataa (SEQ ID NO: 35) aacttgatggcrttkccaacmaggtt (SEQ ID NO: 36) ctctgtgaacttgatrgcrttkccaac (SEQ ID NO: 37) atacacttttcncggtcacccatcat (SEQ ID NO: 38) tccatctgbgcctggatacacttttc (SEQ ID NO: 39) ggcttvgavagatactcgtccatctg (SEQ ID NO: 40).

BIOTECHNOLOGY - Preferred Transformed Cell: The polynucleotide is a polynucleotide complementing the deficiency in hybridsensor kinase in the cell deficient in at least one hybrid-sensor kinase in which the polynucleotide has been introduced. The cell is a microorganism, particularly budding yeast. The osmosensing histidine kinase having no transmembrane region is an osmosensing histidine kinase having no transmembrane region and having a mutation which confers resistance to any of a dicarboxylimide antifungal compound, an aromatic hydrocarbon antifungal compound and a phenylpyrrole antifungal compound to the cell. The osmosensing histidine kinase is derived from a plant-pathogenic filamentous fungus and has no transmembrane region. The osmosensing histidine kinase has an amino acid sequence of 1315, 1307, 1293, 1307, 1438 or 124 amino acids (SEQ ID NO: 1, 16, 41, 55, 68 or 90, respectively) given in the specification. The nucleotide sequence encoding an amino acid sequence of the osmosensing histidine kinase is a sequence of 3948, 3924, 3882, 3924, or 4317 bp (SEQ ID NO: 2, 17, 42, 56, or 69, respectively) also given in the specification. Preferred Method: Assaying the antifungal activity of a substance comprises culturing a transformed cell defined above in the presence of a test substance, measuring an amount of intracellular signal transduction from the osmosensing histidine kinase having no transmembrane region expressed in the cultured transformed cell or an index value having the correlation, and assessing the antifungal activity of the test substance based on a

difference between an amount of intracellular signal transduction or an index value having the correlation measured and a control. The amount of intracellular signal transduction or the index value having the correlation is an amount of growth of the transformed cell. Identifying an antifungal compound comprises selecting an antifungal compound based on the antifungal activity assessed in the assaying method defined above. Killing a fungus comprises identifying an antifungal compound by the method above and contacting the fungus with the identified antifungal compound. Obtaining the polynucleotide above comprises amplifying a desired polynucleotide by PCR using the oligonucleotide above and recovering the amplified desired polynucleotide.

ACTIVITY - Fungicide. No biological data given.

MECHANISM OF ACTION - None given.

USE - The transformed cell is useful for assaying the antifungal activity of a substance and identifying an antifungal compound which is useful for killing a fungus (claimed).

EXAMPLE - BcOS-1 DNA was cloned into a shuttle vector p415ADH (ATCC87312) replicable in yeast and Escherichia coli. Both were digested, separated by agarose gel electrophoresis, and a part of the gel containing a desired DNA was excised. The BcOS-1 DNA was inserted between SpeI and Pstl sites in the multicloning site of the shuttle vector. A nucleotide sequence of the resulting expression plasmid was analyzed after a sequencing reaction. The nucleotide sequence of 3948 bp (SEQ ID NO: 2) given in the specification was obtained and it was confirmed that the expression plasmid pADHBcOS1 harbored a DNA having a nucleotide sequence encoding an amino acid sequence of BcOS-1. The prepared expression plasmid was introduced into each of budding yeast (Saccharomyces cerevisiae AH22 strain) (IFO10144) and TM182 strain. By utilizing disappearance of leucine auxotrophy in the resulting transformed budding yeast , the transformed budding yeast AH22 strain was selected on a Glu-Leu agar medium, and the transformed budding yeast TM182 was selected on a Gal-Ura-Leu agar medium. It was confirmed that the resulting TM182-BcOS1 grows even when transplanted to a Glu-Ura-Leu medium. (211 pages)

L12 ANSWER 4 OF 4 MEDLINE on STN DUPLICATE 2

ACCESSION NUMBER: 2001076412 MEDLINE DOCUMENT NUMBER: PubMed ID: 11073911

TITLE: Novel role for an HPt domain in stabilizing the

phosphorylated state of a response regulator domain.

AUTHOR: Janiak-Spens F; Sparling D P; West A H

CORPORATE SOURCE: Department of Chemistry and Biochemistry, University of

Oklahoma, Norman; Oklahoma 73019, USA.

CONTRACT NUMBER: GM59311 (NIGMS)

SOURCE: Journal of bacteriology, (2000 Dec) Vol. 182, No. 23, pp.

6673-8.

Journal code: 2985120R. ISSN: 0021-9193.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

(RESEARCH SUPPORT, U.S. GOV'T, NON-P.H.S.)

(RESEARCH SUPPORT, U.S. GOV'T, P.H.S.)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: · 200101

ENTRY DATE: Entered STN: 22 Mar 2001

Last Updated on STN: 14 Feb 2003 Entered Medline: 11 Jan 2001

AB Two-component regulatory systems that utilize a multistep phosphorelay mechanism often involve a histidine-containing phosphotransfer (HPt) domain. These HPt domains serve an essential role as histidine-phosphorylated protein intermediates during phosphoryl transfer from one response regulator domain to another. In Saccharomyces cerevisiae, the YPD1 protein facilitates phosphoryl transfer from

a hybrid sensor kinase, SLN1, to two distinct response regulator proteins, SSK1 and SKN7. Because the phosphorylation state largely determines the functional state of response regulator proteins, we have carried out a comparative study of the phosphorylated lifetimes of the three response regulator domains associated with SLN1, SSK1, and SKN7 (R1, R2, and R3, respectively). The isolated regulatory domains exhibited phosphorylated lifetimes within the range previously observed for other response regulator domains (i.e., several minutes to several hours). However, in the presence of YPD1, we found that the half-life of phosphorylated SSK1-R2 was dramatically extended (almost 200-fold longer than in the absence of YPD1). This stabilization effect was specific for SSK1-R2 and was not observed for SLN1-R1 or SKN7-R3. Our findings suggest a mechanism by which SSK1 is maintained in its phosphorylated state under normal physiological conditions and demonstrate an unprecedented regulatory role for an HPt domain in a phosphorelay signaling system.

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EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
L1	1230	budding adj yeast\$2	US-PGPUB; USPAT	OR	OFF	2008/01/18 09:36
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L3	31129	l1 or l2	US-PGPUB; USPAT	OR	OFF	2008/01/18 09:37
L4	135	l3 adj (transform\$3 or tranfect\$3)	US-PGPUB; USPAT	OR	OFF	2008/01/18 09:37
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L11	3	I4 and I10	US-PGPUB; USPAT	OR	OFF	2008/01/18 09:39

US 20050272924 A1 US-PGPUB US 20040171695 A1 US-PGPUB US 20040137594 A1 US-PGPUB US 6803191 B2 USPAT US 6768041 B2 USPAT US 6716625 B1 USPAT US 6673777 B1 USPAT US 6518021 B1 USPAT US 6495356 B1 USPAT US 6433154 B1 USPAT US 6359198 B1 USPAT US 6143728 A USPAT US 5939306 A USPAT

US 20070185314 A1 US-PGPUB
US 20070141578 A1 US-PGPUB
US 20060269951 A1 US-PGPUB
US 20060057607 A1 US-PGPUB
US 20040137594 A1 US-PGPUB
US 20030175930 A1 US-PGPUB
US 20030165932 A1 US-PGPUB
US 20030023032 A1 US-PGPUB
US 7208612 B2 USPAT
US 7183099 B2 USPAT

	Document ID	Kind Codes	Source	Issue Date	Pages
11	US 20070248967 A1		US- PGPUB	20071025	101
2	US 20070134763 A1	l	US - PGPUB	20070614	77
13	US 20040137594 Al	1	US- PGPUB	20040715	37

	Title
1	Reporter Assay Using Secrectory Luminescent Enzymes
2	Anti-virus therapy for respiratory diseases
3	Transformed cell with enhanced sensitivity to antifungal compound and use thereof